



# **STIC Search Report**

*Rush*

## **Biotech-Chem Library**

**STIC Database Tracking Number: 124843**

**TO: Christian Fronda  
Location: REM-3B61/3C70  
Art Unit: 1652  
Friday, June 18, 2004**

**Case Serial Number: 10/023889**

**From: Mary Jane Ruhl  
Location: Biotech-Chem Library  
Remsen 1-A-62  
Phone: 571-272-2524**

**maryjane.ruhl@uspto.gov**

### **Search Notes**

Examiner Fronda,

Here are the results for your recent search request.

Please feel free to contact me if you have any questions about these results.

Thank you for using STIC services. We appreciate the opportunity to serve you.

Sincerely,

Mary Jane Ruhl  
Technical Information Specialist  
STIC  
Remsen 1-A-62  
Ext. 22524

124843  
**STIC-Biotech/ChemLib**

**From:** Chan, Christina  
**Sent:** Wednesday, June 16, 2004 5:38 PM  
**To:** Fronda, Christian; STIC-Biotech/ChemLib  
**Subject:** RE: Rush Search for Serial No. 10/023,889

Please rush. Thanks Chris

*Chris Chan*

TC 1600 New Hire Training Coordinator and SPE 1644 & 1642  
(571)-272-0841  
Remsen, 3E89

-----Original Message-----

**From:** Fronda, Christian  
**Sent:** Wednesday, June 16, 2004 9:18 AM  
**To:** Chan, Christina  
**Subject:** Rush Search for Serial No. 10/023,889  
**Importance:** High

I would like to request a Rush Search for Serial No. 10/023,889 as listed below since it is an overdue date case.  
Thank you.

Christian L. Fronda  
Art Unit 1652  
Office REM B61  
Mailbox REM C70  
(571)272-0929

Please perform sequence search and interference search for Serial No. 10/023,889

1. Please search SEQ ID Nos.:1-7, 17, and 18 against amino acid commercial, PBPub, and issued databases.

**Please save on COMPUTER DISKETTES.**

Thank you very much.

Christian Fronda  
Art Unit 1652  
Mailbox REM 3C70  
Office REM 3B61  
(517)272-0929

Searcher: \_\_\_\_\_  
Phone: \_\_\_\_\_  
Location: \_\_\_\_\_  
Date Picked Up: \_\_\_\_\_  
Date Completed: \_\_\_\_\_  
Searcher Prep/Review: \_\_\_\_\_  
Clerical: \_\_\_\_\_  
Online time: \_\_\_\_\_

TYPE OF SEARCH:  
NA Sequences: \_\_\_\_\_  
AA Sequences: \_\_\_\_\_  
Structures: \_\_\_\_\_  
Bibliographic: \_\_\_\_\_  
Litigation: \_\_\_\_\_  
Full text: \_\_\_\_\_  
Patent Family: \_\_\_\_\_  
Other: \_\_\_\_\_

VENDOR/COST (where applic.)  
STN: \_\_\_\_\_  
DIALOG: \_\_\_\_\_  
Questel/Orbit: \_\_\_\_\_  
DRLink: \_\_\_\_\_  
Lexis/Nexis: \_\_\_\_\_  
Sequence Sys.: \_\_\_\_\_  
WWW/Internet: \_\_\_\_\_  
Other (specify): \_\_\_\_\_

Pending Nucleic Acid and Pending Amino Acid database searches generate two sets of results each. The Pending databases have been split into two pairs to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions `.rapm` and `.rapn`

Searches run against the Amino Acid Pending database produce two sets of results, with the extensions `.rapm` and `.rapn`

*Because they contain data that is confidential, the results of Pending database searches should not be left in the case.*